SEQUENCE LISTING

5	<110> Green, Allan Singh, Surinder Liu, Qing	
	<120> METHOD OF MODIFYING THE CONTENT OF COTTON SEED OIL	
10	<130> p:\oper\mro\specifications\cottonseed oil US	
<u>).</u> U	<140> US <141> 2001-04-18	
15	<150> US 60/198,124 <151> 2000-04-18	
	<160> 35	
20	<170> PatentIn version 3.0	
	<210> 1 <211> 1493	
25	<212> DNA <213> cotton	
	<220> <221> CDS <222> (13)(1200)	
30	<400> 1	
	cgaaaagaaa aa atg got ttg aat ttt aat goc atc goc tog aaa tot cag Met Ala Leu Asn Phe Asn Ala Ile Ala Ser Lys Ser Gln 1 5 10	51
35	aag ete eet tge tit get ett eea eea aag gee aee ett aga tet eee	99
	Lys Leu Pro Cys Phe Ala Leu Pro Pro Lys Ala Thr Leu Arg Ser Pro 15 20 25	
40	aag ttt tee atg ate tee ace att eet tet gge tee aaa gag gtt ggg Lys Phe Ser Met Ile Ser Thr Ile Pro Ser Gly Ser Lys Glu Val Gly 30 35 40 45	147
4.5	aat ctg aaa aag cet tte acg cet cea aag gag gtg cet gtt eag ate	195
45	Asn Leu Lys Lys Pro Phe Thr Pro Pro Lys Glu Val Pro Val Gln Ile 50 55 60	
50	acc cac too atg cog cot cac aag att gag atc ttt aaa tot ttg gag Thr His Ser Met Pro Pro His Lys Ile Glu Ile Phe Lys Ser Leu Glu 65 75	243
50	,,,	
	ggo tgg got gag aac aac att otg act cac ctc aaa cca gtt gag aaa	291
	ggc tgg gct gag aac aac att ctg act cac ctc aaa cca gtt gag aaa Gly Trp Ala Glu Asn Asn Ile Leu Thr His Leu Lys Pro Val Glu Lys 80 85 90	291
55	6ly Trp Ala Glu Ash Ash Ile Leu Thr His Leu Lys Pro Val Glu Lys 80 85 90 tgt tgg caa ccc gcc gac ttt ctt cca gat cct aat tct gat gga ttt	291 339
55	Gly Trp Ala Glu Asn Asn Ile Leu Thr His Leu Lys Pro Val Glu Lys	
55 60	tgt tgg caa ccc gcc gac ttt ctt cca gat cct aat tct gat gga ttt Cys Trp Gln Pro Ala Asp Phe Leu Pro Asp Pro Asn Ser Asp Gly Phe	

	gat	tac	ttt	gta	gtt	ttg	gtt	ggt	gat	atq	- atc	acc	gag	gaa	gee	ctt	435
							Val		_	-				-	-		
5						-	ctt Leu			_	-			-			483
10							Pro		-								531
15	-	-	-				ggt Gly 180	_	_			_				-	57 9
20					_	-	agg Arg							_		_	627
			-		_	_	ect Pro					_			_		675
25					-		caa Gln	_		_							723
30			_		-		aag L y s				_			_	_		771
35							tca Ser 260				-				_		819
40	Thr 270	Lys	Ile	Val	Glu	Lys 275	ctg Leu	Phe	Glu	Ile	Авр 280	Pro	Aap	Glu	Thx	Val 285	867
							atg Met										915
45	ttc Phe	atc Ile	tat Tyr	gat Asp 305	Gly	aga Arg	gat Asp	tat Tyr	Asn	Leu	ttt Phe	gac Asp	His	tac Tyr 315	tca Ser	get Ala	963
50	gtt Val	Ala	caa Gln 320	aga Arg	atc Ile	ej aaa	gtt Val	tac Tyr 325	act Thr	get Ala	aag Lys	gac Asp	tat Tyr 330	gtt Val	gat Asp	ata Ile	1011
55	gta Val	gag Glu 335	cac His	ctg Leu	gtg Val	gat Asp	ega Arg 340	tgg Trp	aag Lys	gtg Val	aag Lys	gag Glu 345	cta Leu	gct Ala	e1Å ååå	ctt Leu	1059
60	tca Ser 350	gcc Ala	gag Glu	ejā āāā	cgt Arg	aaa Lys 355	gct Ala	cag Gln	yab ggc	tac Tyr	ttg Leu 360	tgt Cys	tca Ser	ctt Leu	cct Pro	tcg Ser 365	1107
	aga Arg	att Ile	aga Arg	agg A rg	tta Leu 370	gag Glu	gag Glu	aga Arg	gcg Ala	caa Gln 375	gaa Glu	aag Lys	Ala	aag Lys	gaa Glu 380	gca Ala	1155

ج					gaa gtg aaa ctt Glu Val Lys Leu 395	1200
5	taggtcatç	ga aatacagt	ta agactec	tgc aatgcatttg	aggaaacaaa cacga	agaag 1260
	aattgcgtg	gg ctttggtt	ag ggtagca	cat gttttctgga	tgtgttgtgt cctta	aaaaa 1320
10	taatgccga	at agcggcag	ct gtgatag	ttt tagatgtttg	ttttcataat gtctg	ttata 1380
	togttgtac	eg agtagtat	gt gttgttt	ttg ttgaaacaat	ottcatatct tagtg	ataaa 1440
15	tgataatgo	ct gtgtagtc	at agttttt	a gt tigcaalaaa	aaaaaaaaa aaa	1493
20	<210> 2 <211> 39 <212> PR <213> co					
	<400> 2					
25	Met Ala I	Leu Asn Phe 5	Asn Ala I	le Ala Ser Lys 10	Ser Gln Lys Leu 15	Pro
30	Cys Phe A	Ala Leu Pro 20	Pro Lys A	la Thr Leu Arg 25	Ser Pro Lys Phe 30	Ser
		Ser Thr Ile 35	Pro Ser G		Val Gly Asn Leu 45	Lys
35	Lys Pro P 50	Phe Thr Pro	Pro Lys G	lu Val Pro Val	Gln Ile Thr His	Ser
40	Met Pro P 65	Pro His Lys	Ile Glu I 70	le Phe Lys Ser 75	Leu Glu Gly Trp	Ala 80
45	Glu Asn A	Asn Ile Leu 85	Thr His L	au Lys Pro Val 90	Glu Lys Cys Trp 95	Gln
50	Pro Ala A	lsp Phe Leu 100		ro Asn Ser Asp 105	Gly Phe His Glu 110	Gln
	_	Slu Leu Arg 115	_	la Lys Glu Ile 20	Pro Asp Asp Tyr 125	Phe
55	Val Val I	ieu Val Gly	Asp Met I 135	le Thr Glu Glu	Ala Leu Ser Thr 140	Tyr
60	Gln Thr M	Met Leu Asn	Thr Leu A	sp Gly Thr Arg 155	Asp Glu Thr Gly	Ala 160

	Ser	Leu	Thr	Pro	Trp 165	Ala	Ile	Trp	Thr	Arg 170	Ala	Trp	Thr	Ala	Glu 1 75	Glu
5	Asn	Arg	His	Gly 180	Asp	Leu	Leu	Asn	Lys 185	Tyr	Leu	ΊΫ́Σ	Leu	ser 190	Gly	Arg
10	Val	Asp	Met 195	Arg	Gln	Ile	Glu	Arg 200	Thr	Ile	Gln	Tyr	Leu 205	Ile	Gly	Ser
15	Gly	Met 210	Asp	Pro	His	Thr	Glu 215	Asn	Ser	Pro	Туг	Arg 220	GlĀ	Phe	Ile	Tyr
	Thr 225	Ser	Phe	Gln	Glu	Arg 230	Ala	Thr	Phe	Ile	Ser 235	His	Gly	Asn	Thr	Gly 240
20	Arg	Leu	Ala	Lys	Glu 245	Tyr	Gly	Авр	Ile	Asn 250	Leu	Ala	Gln	Ile	Cys 255	Cly
25	Ser	Ile	Ala	Ser 260	Asp	Glu	Lys	Arg	His 265	Glu	Thr	Ala	Tyr	Thr 270	Lys	Ile
30	Val	Glu	Lys 275	Leu	Phe	Glu	Ile	Asp 280	Pro	Аар	Glu	Thr	Val 285	Leu	Ala	Phe
35	Ala	Asp 290		Met	Lys	Lys	Lys 295	Ile	Ala	Met	Pro	Ala 300	Glu	Phe	Ile	Tyr
	Asp 305	Gly	Arg	Asp	Tyr	Asn 310	Leu	Phe	Asp	His	Tyr 315	Ser	Ala	Val	Ala	Gln 320
40	Arg	Ile	Gly	Val	Tyr 325	Thr	Ala	Lys	Asp	Tyr 330	Val	Asp	Ile	Va1	61u 335	His
45	Leu	Val	Asp	Arg 340	Trp	Lys	Val	Lys	Glu 345	Leu	Ala	Gly	Leu	Ser 350	Ala	Glu
50	ely	Arg	Lys 355	Ala	Gln	qaA	Туг	Leu 360	Суз	Ser	Leu	Pro	Ser 365	Arg	Ile	Arg
55	Arg	Leu 370	Glu	Glu	Arg	Ala	Gln 375	Glu	Lys	Ala	Lys	Glu 380	Ala	Pro	Ser	Val
	Pro 385	Phe	Ser	Trp	Ile	Phe 390	Asp	Arg	Glu	Val	Lys 395	Leu				
60	<210 <211 <212	L>]	3 1411 DNA													

<213> cotton

<220>

<221> CDS <222> (79)..(1233)

	<400>	3	acca	acac	gc c	ttet	ttgc	c te	gtgt	ttca	tca	eetg	geg '	ttaa	actgct	60
10	ttottt	aag	ccag	caaa	-	ggt Gly	-			-				_		111
15	ata aag Ile Lys															159
20	eet eeg Pro Pro		_			-		-		_					-	207
25	ttt cgc Phe Arg 45	-				_						_		-		255
30	tgc tta Cys Lev 60	-							_							303
	ctc cca Leu Pro															351
35	caa ggt						_		-		_					399
40	cac cac															447
45	atc ctt Ile Leu 125	His	Ser	Ala	Leu	Leu 130	Val	Pro	Tyr	Phe	Ser 135	Trp	ГÀЗ	Ile	Ser	495
50	cac cgc His Arg 140	Arg	His	His	Ser 145	Asn	Thr	Gly	Ser	Met 150	Glu	Arg	Asp	Glu	Val 155	543
	ttc gtg Phe Val	Pro	aaa Lys	Pro 160	aag Lys	tet Ser	aaa Lys	tta Leu	tca Ser 165	tgc Cys	ttt Phe	Al a	rys 888	TYF 170	tta Leu	591
55	aac aat Asn Asn	Pro	920 175	Gly ggt	cga Arg	gtt Val	cta Leu	tct Ser 180	ctt Leu	gta Val	gtc Val	aca Thr	ttg Leu 185	act Thr	ctt Leu	639
60	ggt tgg Gly Trp	Pro 190	atg Met	tac Tyr	tta Leu	gcc Ala	ttc Phe 195	aac Asn	gtt Val	tcg Ser	Gly	cga Arg 200	tac Tyr	tat Tyr	gat Asp	687
	cga tta	gct	tcc	cac	tat	aac	cct	tat	ggc	ccc	att	tac	tec	gat	cgc	735

	Arg	Leu 205	Ala	Ser	His	Tyr	Asn 210	Pro	Tyr	Gly	Pro	Ile 215	Tyr	Ser	Авр	Arg	
	gag	agg	cta	caa	att	tac	atc	tcc	gat	act	aat	ata	ttt	aca	ota	att	783
5		Arg															
_	220					225					230					235	
	tat	gta	ctt	tat	aaq	att	act	σca	aca	aaa	aaa	cta	act	taa	CEE	tta	831
		Val															
10					240					245					250		
	tgc	act	tat	ggg	gtg	cct	cta	ctt	att	gtg	aat	gcc	ttc	ctt	gtg	ttg	879
	Сув	The	Tyz		Val	Pro	Leu	Leu		Val	Asn	Ala	Phe		Val	Leu	
15				255					260					265			
13	atc	acc	tac	ttg	caa	cat	act	cac	tcg	gca	ttg	ccg	cat	tat	gac	teg	927
	Ile	Thr		Lou	eju	His	The		Ser	Ala	Leu	Pro		Tyr	Asp	Ser	
			270					275	•				280				
20		gaa															975
	Ser	Glu 285	Trp	Asp	Trp	Leu		Gly	Ala	Leu	Ser		Met	Asp	Arg	Asp	
		203					290					295					
		ggg															1023
25	Phe 300	СŢĀ	Val	Leu	Asn	Lу в 305	Val	Phe	His	Asn	Ile 310	Thr	Авр	Thr	His		
	500					303					310	•				315	
		cat															1071
30	ATA	His	HŢS	Leu	2he 320	Ser	Thr	Met	Pro	His 325	TYE	His	Ala	Met	Glu 330	Ala	
30					720					323					330		
		aaa															1119
	THE	Lys	ATG	335	гĀЗ	Pro	ITe	Leu	GLY 340	Lys	TYT	TYT	Pro	Phe 345	Asp	Gly	
35									_								
	aca aca	ecg Pro	att	tac	aag	gca	atg	tgg m	agg	gag	gca	aaa	gag	tgc	ctt	tac	1167
	****	220	350	+YL	LYS	VT4	mec	355	wrd	GIU	WIG	тАа	360	Cys	Ten	TYT	
40	gtt Val	gag Glu	Pro	gac	gtt Val	ggt.	ggt	ggt Glv	ggt	ggt	ggt	agc	aaa Lua	ggt	gtt	ttt	1215
		365					370	-1	413	411	GLY	375	-y -	GTĀ	441	2110	
	t .c.c						4						.				
45		tat Tyr					taaa	gaco	ga c	CEBS	rgç	et ge	tago	tggo	2		1263
	380	_	_		_	385											
	caac	TAAP	tc s	acat	'aaa=		-20+6	· = + + =		*						taatt	1323
	-99-	,,				ic gi	-400		. yac	· cay (ع ي د د	dact	.ayyy	aa ç	LUA	icaact	. 1323
50	aato	ıgtaç	ıga a	ıaatç	rtgga	a ta	igttg	rccta	gta	igtti	tat	gtat	taaç	rtg t	tgt	ttaat	1383
	aaac	tata	tg c	jtags	aaaa	18 8E	sasa	aa									1411
				-													
55	<210	> 4	ı														
55	<211		85														
	<212	_	RT														
	<213	s> c	otto	'n													
60	<400	> 4	l														
	Met	GJĀ	Ala	Glv	Glv	Ara	Met	Pro	Tle	Apr	GI w	Tle	Tare	G7 **	G) **) an	
	1	-			5					10	1		-3-		15		

5	Arg	Cly	Ser	Val 20	Asn	Arg	Val	Pro	11e 25	Glu	Lys	Pro	Pro	Phe 30	Thr	Leu
	Gly	Gln	Ile 35	Lys	Gln	Ala	Ile	Pro 40	Pro	His	СХа	Phe	Arg 45	Arg	Ser	Leu
10	Leu	Arg 50	Ser	Phe	Ser	Tyr	Val 55	Val	His	Asp	Leu	С у в 60	Leu	Ala	Ser	Phe
15	Phe 65	Tyr	Tyr	Ile	Ala	Thr 70	Ser	Туг	Phe	His	Phe 75	Leu	Pro	Gln		Phe 80
20	Ser	Tyr	Ile	Ala	Trp 85	Pro	Val	Tyr	Trp	Val 90	Leu	Gln	Gly	Сув	Ile 95	Leu
25	Thr	Gly	Val	Т <u>тр</u> 100	Val	Ile	Ala	His	Glu 105	Trp	Gly	His	His	Ala 110	Phe	Arg
	Asp	Tyr	Gln 115	Trp	Val	Asp	Asp	Thr 120	V al	Gly	Leu	Ile	Leu 125	His	Ser	Ala
30	Leu	Leu 130	Val	Pro	Tyr	Phe	Ser 135	Trp	Lys	Ile	Ser	His 140	Arg	Arg	His	His
35	Ser 145	Asn	Thz	Cly	Ser	Met 150	Glu	Arg	Asp	Glu	Val 155	Phe	V al	Pro	Lys	Pro 160
40	ГÀв	Ser	Lys	Leu	Ser 165	Cys	Phe	Ala	Lys	Tyr 170	Leu	Asn	Asn	Pro	Pro 175	Gly
45	Arg	Val	Leu	Ser 180	Leu	Val	Val	Thr	Leu 185	Thr	Leu	Gly	Trp	Pro 190	Met	Tyr
	Leu	Ala	Phe 195	Asn	Val	Ser	Gly	Arg 200	Tyr	Tyz	Авр	Arg	Leu 205	Ala	Ser	His
50	Tyr	Asn 210	Pro	Туг	Gly	Pro	Ile 215	Tyr	Ser	qaA	Arg	Glu 220	Arg	Leu	Gln	Val
55	ту г 225	Ile	Ser	Asp	The	61 ₃ 230	Ilə	Phe	Ala	Val	11e 235	Tyr	Val	Leu	Tyr	Lуа 240
60	Ile	Ala	Ala	Thr	Lys 245	Gly	Leu	Ala	Trp	Leu 250	Leu	Сув	The	Туг	Gly 255	Val

Pro Leu Leu Ile Val Asn Ala Phe Leu Val Leu Ile Thr Tyr Leu Gln

5	His Thr His Ser Ala Leu Pro His Tyr Asp Ser Ser Glu Trp Asp Trp 275 280 285	
10	Leu Arg Gly Ala Leu Ser Thr Met Asp Arg Asp Phe Gly Val Leu Asn 290 295 300	
	Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Leu Phe 305 310 315 320	
15	Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile Lys 325 330 335	
20	Pro Ile Leu Gly Lys Tyr Tyr Pro Phe Asp Gly Thr Pro Ile Tyr Lys 340 345 350	
25	Ala Met Trp Arg Glu Ala Lys Glu Cys Leu Tyr Val Glu Pro Asp Val 355 360 365	
30	Gly Gly Gly Gly Gly Ser Lys Gly Val Phe Trp Tyr Arg Asn Lys 370 375 380	
	Phe 385	
35	<210> 5 <211> 1422 <212> DNA <213> cotton	
40	<220> <221> CDS <222> (98)(1246)	
45	<400> 5 taaaaaaaa aggcatttot ttoatottaa agagacagog aggaagocao gaagataata	60
50	gagtgatttt caatotocat tttaagggtg tggaaca atg ggt gct gga ggc aga Met Gly Ala Gly Gly Arg 1	115
50	atg tog gtt cca acg agt cca aaa aaa ccc gaa ttc aac tca ctg aag Met Ser Val Pro Thr Ser Pro Lys Lys Pro Glu Phe Asn Ser Leu Lys 10 15 20	163
55	cga gtt cca tac tca aag cca ccc ttc act ctg agt gaa atc aag aaa Arg Val Pro Tyr Ser Lys Pro Pro Phe Thr Leu Ser Glu Ile Lys Lys 25 30 35	211
60	gcc atc cca cca cac tgt ttc cag cgc tcc gtt tta cgc tca ttc tca Ala Ile Pro Pro His Cys Phe Gln Arg Ser Val Leu Arg Ser Phe Ser 40 45 50	259

		ctc Leu															:	307
5		aat Asn															:	355
10		ctt Leu																403
15		gcc Ala							_		-	-					•	451
20	-	gac Asp 120												-			4	199
		tct Ser				_			-								5	547
25		gaa Glu		-	-			-		_						_	5	595
30		tgg Trp	-							_				_			(643
35		att Ile							-								6	591
40		ggc Gly 200				-			-	_			-				7	739
		ata Ile	_		-		_	_							-	_	Ī	787
45		gtc Val															E	935
50		gta Val	-		_		-	_			_			_		-	ŧ	983
55		gcc Ala															!	931
60		ccg Pro 280															!	979
-	tca Ser 295	act Thr	gtg Val	gac Asp	aga Arg	gat Asp 300	tat Tyr	ggg ggg	att Ile	tta Leu	aac Asn 305	aag Lys	gtt Val	ttc Phe	cat His	aac Asn 310	1	027

_	ata acc gac act cat gtg gct cat cat ttg ttt tcg aca atg cct cac Ile Thr Asp Thr His Val Ala His His Leu Phe Ser Thr Met Pro His 315 320 325	1075
5	tat cat geo atg gtg gcc acc aag gcg ata aag ccc ata ttg ggg gaa Tyr His Ala Met Val Ala Thr Lys Ala Ile Lys Pro Ile Leu Gly Glu 330 335 340	1123
10	tac tat cag ttc gat ggg atg cet gtc tat aag gcg ata tgg agg gag Tyr Tyr Gln Phe Asp Gly Met Pro Val Tyr Lys Ala Ile Trp Arg Glu 345 350 355	1171
15	gcg aag gag tgt ctc tac gtt gaa cca gat gag ggc gac aag gat aaa Ala Lys Glu Cys Leu Tyr Val Glu Pro Asp Glu Gly Asp Lys Asp Lys 360 365 370	1219
20	ggt gtg ttt tgg ttt aga amc amg ctt tmamatatttg cattitacct Gly Val Phe Trp Phe Arg Asn Lys Leu 375 380	1266
	taggeatgtt ctagtegttg atgttttaag gatattttag eegacataet tggtttteet	1326
25	ttttgggact ttttagcttt gtatttgcag acaataatct tgttcactat taaataatgg	1386
4 3	tagaaataaa tacacagcat ggattggcaa taaaaa	1422
30	<210> 6 <211> 383 <212> PRT <213> cotton	
	<400> 6	
35	Met Gly Ala Gly Gly Arg Met Ser Val Pro Thr Ser Pro Lys Lys Pro 1 5 10	
40	Glu Phe Asn Ser Leu Lys Arg Val Pro Tyr Ser Lys Pro Pro Phe Thr 20 25 30	
45	Leu Ser Glu Ile Lys Lys Ala Ile Pro Pro His Cys Phe Gln Arg Ser 35 40 45	
50	Val Leu Arg Ser Phe Ser Tyr Leu Leu Tyr Asp Phe Ile Leu Ala Ser 50 55 60	
	Leu Phe Tyr His Val Ala Thr Asn Tyr Phe Pro Asn Leu Pro Gln Ala 65 70 75 80	
5 <i>5</i>	Leu Ser Asn Val Ala Trp Pro Lou Tyr Trp Ala Met Gln Gly Cys Ile 85 90 95	
60	Leu Thr Gly Val Trp Val Ile Ala His Glu Cys Gly His His Ala Phe	

15

Ser	Asp	Tyr	Gln	Trp	Leu	Asp	Asp	Thr	Val	Gly	Leu	Ile	Leu	His	Şe≭
		115					120					125			

- 5 Ser Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser His Arg Arg His 130 135 140
- His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys
 10 145 150 155
- Lys Lys Ser Gly Leu Arg Trp Trp Ala Lys His Phe Asn Asn Pro Pro 165 170 175
- Gly Arg Phe Leu Ser Ile Thr Ile Gln Leu Thr Leu Gly Trp Pro Leu 180 185 190
- Tyr Leu Ala Phe Asn Val Ala Gly Arg Fro Tyr Asp Arg Phe Ala Cys
 195 200 205
- 25 His Tyr Asp Pro Tyr Gly Pro Ile Phe Ser Asp Arg Glu Arg Leu Gln 210 215 220
- Ile Tyr Ile Ser Asp Ala Gly Val Leu Ala Val Ala Tyr Ala Leu Tyr 30 225 230 230
- Arg Leu Val Leu Ala Lys Gly Val Gly Trp Val Ile Ser Val Tyr Gly 245 250 255
 - Val Pro Leu Leu Val Val Asn Ala Phe Leu Val Met Ile Thr Tyr Leu 260 265 270
- Gln His Thr His Pro Ser Leu Pro His Tyr Asp Ser Ser Glu Trp Asp
 275 280 285
- 45 Trp Met Arg Gly Ala Leu Ser Thr Val Asp Arg Asp Tyr Gly Ile Leu 290 295 300
- Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Leu 50 305 310 315 320
- Phe Ser Thr Met Pro His Tyr His Ala Met Val Ala Thr Lys Ala Ile 325 330 335
 - Lys Pro Ile Leu Gly Glu Tyr Tyr Gln Phe Asp Gly Met Pro Val Tyr 340 345 350
- Lys Ala Ile Trp Arg Glu Ala Lys Glu Cys Leu Tyr Val Glu Pro Asp 355 360 365

Glu Gly Asp Lys Asp Lys Gly Val Phe Trp Phe Arg Ash Lys Leu 370 375 380

5	<210> 7 <211> 5000 <212> DNA <213> cott						
15	<220> <221> Int:	35) (5006)					
20	<400> 7	39)(4998) ttttataggc	atttaatagg	gaggaattt	gaaaaatttc	atgaataaag	60
25		actgagaaaa					120
	aaaggcccaa	ctaatccaaa	caaaacttga	gtgttacaat	ctaaccctag	tctggcaacg	180
	gatacgggtt	aagggtgtta	caacctttac	agtgatcaac	gaacaaacct	tgagtggatt	240
30	tggatttgac	cccctaccc	cactacacac	aaggaagaat	gttagtttag	ttattcaata	300
	gctactaagt	tggtttacat	atatatacaa	gttccacact	tgattctcaa	tcaatgtgag	360
	actaatgctt	ttcatttctc	tcaacataat	tcacaagtag	cttactttga	gtatcaattt	420
35	ttcattcatc	actcaatcat	tttgagcata	tgatatattg	ttgtaaatgt	ctaatggagt	480
	agaatataaa	attataattt	tatgattcaa	cttttcacct	ttaccaatag	aaaatatgee	540
40	tcaaagtttt	caaaaaatca	tattttttct	aatagaaata	actttagaca	tcaagaatct	600
	acgaataaaa	tttaaataac	ttttttctcc	aatottogat	acttgctatt	aaattaactt	660
	aaattettet	acttgagatt	ttgatgcatg	cgacaaaaat	tgagattaga	atccatgatg	720
45	tggaaagcaa	aacasstatg	taagcaaata	ttgtcttgtc	gtaaaaccat	ttgattttt	780
		ttcattgaaa					840
50		aacttaaacg	-			-	900
50		gegatactte		-			960
				_			
55		cgtgatggag					1020
	ttattattaa	tattttattg	ataaaagtat	tagaaatttt	tgtattaaaa	gtcagattat	1080
	attttatatt	ttaaaaataa	ataagttagt	cattetgtgt	tagatcaaat	agcaaaacaa	1140
60	tagaaataga	tgaaattttc	aataaaaaag	gaccagttta	ctctttgaac	taacgcacaa	1200
	tgactaattt	accattttta	gtagatgagg	taaaatataa	tctagctcct	tgtacagggg	1260

	cttccgtgat	acttttaccc	tattgtatct	cttctcaacg	ataaatanaa	atatgtttta	1320
	gasaattttg	tttccaaaga	taattacasa	gttaagtcaa	acaagcagta	acattgtttc	1380
5	acttaatttc	cctttcgaaa	gaaaaactct	tatttagaat	aattgtcatt	caaagtaact	1440
	attttttta	gaacagctat	gcttggaaca	atcatgttta	gaacatggct	ccattttaga	1500
	atatggttgt	egtttgagaa	caactcctge	aaaggataac	gaatgtttgg	aacagttett	1560
10	atttagaata	actgcgtttc	aagaataatc	atatttagaa	caacctccat	ttazzacaat	1620
	agtggttttt	tttaasaaga	agagatatta	ttcasastts	gctctttcaa	gaaaagcatc	1680
15	accatagaac	aactattatt	aaaaataaag	ctasattcac	aatttggccc	ttgaagtata	1740
	ctcattttt	gactttggta	tctaaacttt	tetttgeete	aatttgatac	ctaaaatact	1800
20	ctcazattec	attttttgac	agacattaaa	aaaataatct	tatagecaat	cacaaagege	1860
20	cacgtggcgt	ctttatgtaa	aaagaaatat	tttgtttaat	taaatgtata	tacacattaa	1920
	aaaataaaaa	aatatagaac	aacatataaa	ttataaacaa	atctataaaa	atasaattta	1980
25	caaaaataca	ataattgaaa	aaaaattagt	tgaaattaat	aatattatta	aaaaatgtaa	2040
	aacatttgta	asaattataa	aaaagtttta	aaaataattt	totttataaa	attctaaaat	2100
2.0	atataattct	aaaattgtaa	aaaggtatat	asatttcatt	ttttcaatt	actcgaaatt	2160
30	taaatacttt	ttcaattttt	ataacacttt	tttaatttt	atatattatt	ttggatttt	2220
	aaaatttata	attatatatt	tcasaatttt	ataaatttt	ttatatgtta	tatatgattt	2280
35	tttacatttt	taatcastat	aatataaaat	actaaatttt	tttaaaaaaa	tgatacgtgg	2340
	catgttctaa	tategecaca	tgaccggtga	acgctcggtc	aatggccagt	caaagccaaa	2400
40	agtattttt	taatatttaa	ataattaatt	ttatacttct	atttttaatt	ttaatttaat	2460
40	atttttattt	taaataaact	taattgttat	gtgatatttt	ttcactgacc	caaaacatgc	2520
	tatgtggagc	ttttcccatt	agccaaagtt	gccaaataga	tttttcaac	gtgtgttaca	2580
45	aaatggaccg	aaaatagagg	aaattgatac	tttaggtatc	aagttgggat	aaaaaaagtt	2640
	taggtaccaa	agtaaaaaat	ttggtatact	ttaggggcta	aatcataaat	taagootaaa	2700
50	gacaattata	ttgagaataa	attaggttta	gaacgggtca	aaataattct	tattcagtaa	2760
30	caacttttgt	tcaagaacaa	ctcttcaata	acaaaagttt	gtatttcacg	agatgctctt	2820
	attttttgtt	caatatattt	atttgtgcac	tgtaatgatc	gattttaatc	asacataaat	2880
55	aaatgtcttg	attgtgattg	taatttttc	ttagttgtac	aaatatattg	attataattg	2940
	tgatggaaaa	aaataaaaaa	aattaaattt	tagatgaata	aagagtacat	gggctataat	3000
60	tagaattaac	ctasatttgt	ttggttataa	ctagaggttt	tggttcaaag	aattaatttc	3060
80	taaatccgag	tccaacecge	tttggatcag	ccaaaggttc	ttttaaatta	ttttaattat	3120
	ttatatttaa	attaaataat	aaaatatata	aatataatat	aatttcatca	tttattgaat	3180

	cgagttaatc	caaaattgaa	gaatataaac	tcaaactcga	ctcaagatga	ategaacege	3240
_	ttgactaaac	tgacccaacc	caacttgttt	ttgagctaag	tttgagttta	atattttcaa	3300
5	cttcacgttg	gettgaccea	gactgattaa	ttattaaaca	actasasgas	ttttaatata	3360
	catttgaatt	taaatttaaa	taaaaagtat	taatataaga	tctgtgaaaa	gtcgtattta	3420
10	gtgattatat	gatttatgtt	taatttaaat	gttaattatt	tatatttact	tcaacaatag	3480
	tagtaacatt	ctgtaatatg	aatatgaatc	cgatgattga	gaagtgagag	gtgttacata	3540
15	tttactaccg	aggaataect	ttecttccac	gatgaagtcc	ttatogttgc	aacaggtgtt	3600
13	tgggagatat	caaaaagagg	cggggtaatg	atgatgatga	aagcgagaat	catcagaatc	3660
	agaaaaaagg	gaaaagggtg	attaaagaaa	agactctcat	cctcactgat	tetggtettg	3720
20	cttcccaaca	cgtagcatet	aaccataacg	cctcaaatcc	gctgctcctc	tatttatttc	3780
	aaaaccactt	gattaagete	cecataeget	ccataccact	cgcccaaaac	caacacgcct	3840
25	tctttgeete	gtgtttcatc	acctggcgtt	aaactgcttt	ctttaaaggt	acatttctct	3900
23	ttaatttcct	ttttttca	tttcatgttt	ttcatgttaa	tgttgcattg	aagtgataaa	3960
	tttgagtgaa	tgatgtttgg	tatatettet	tagtaactga	ccttttgaaa	atactagcat	4020
30	tttttttaat	atcaagtgaa	agaagaagaa	gaatttcgcc	atgcaaaagc	tttttaaggc	4080
	ttttcttt	ccttagatca	aaatttattt	gtttacttat	actgttcttt	taagcccgaa	4140
35	gaaagaagcc	atggtttcaa	tttttgagag	ttttaaatcc	caaataccag	agagetteat	4200
رر	cgtttattca	tatattttta	aacattttt	aaagcaagaa	cttgtgattt	gtttttaata	4260
	aaatatgcaa	taaatttta	tatttttcgt	aaatttaaaa	tttaattttt	ctacttttaa	4320
40	aatttaaaaa	agtaaatttt	aaaatatacc	tttcattaaa	ttaaattatt	ataagtaatt	4380
	gagtatttt	aattttaaaa	tttcacacat	caaattaaaa	aaaaagttaa	cacttgcact	4440
45	tgattttgaa	aagtaaaagg	attaaatttc	aaattttcag	taaaaggact	aaatttcaaa	4500
,,,	tttttaaaga	gtatagagac	tcctctacat	tttagatttt	aaaatttaaa	tctaacagtt	4560
	aacactttct	taattacttt	acgataaatt	taactaaaaa	attacaatat	taatggttaa	4620
50	aattaaattt	tgaaaagtat	aaagattaaa	ttgtaaattt	tcassasgca	taggaagtta	4680
	tagtatattt	taacctttat	ttattttata	tctggtgagg	ttectgeatg	caccgaagat	4740
55	gtcacctttt	gccagtattt	tecagtgget	tgtttctctc	aaaactacct	tgaatcttga	4800
	gacagaatta	aatatatttt	tggcatttta	ttcattttct	ctctctctat	tttcttttaa	4860
	aaattgettt	agagaattca	gaaaaaatac	tttccaacac	gaaaatttct	tcaaatttat	4920
60	tgtttatatc	taataaatgg	ttgcttaatt	ttggaaaaca	aaagttattg	tagttagttt	4980
	tgcttcttgc	gtgtecagec	agcaaa				5006

	<210> <211>	8 20			
	<212>				
5	<213>	synthetic primer			
	<400>	8			
	atggck	ctsa rgctbcatsc			20
10	<210>	9			
	<211>	20			
	<212>			•	
	<213>	synthetic primer			
15					
	<400>	9		• •	
	teasag	yttm acytgyctat			20
20	<210>	10			
20	<211>	20			
	<212>				
	<213>	synthetic primer			
25	<400>				
	gcatag	gtca tggaccacgt			20
	,		•		
	<210>	11			
30	<211>	17			
	<212>	DNA			
	<213>	synthetic primer			
	<400>				
26	<400>	11 cgac ggccagt			17
35	y caaaa	cgac ggccage			
	<210>	12			
	<211>	19			
40	<212>				
	<213>	synthetic primer			
	<400>	12			
		agct atgaccatg	•		19
45		•			
	<210>	13			
	<211>	8			
50	<212>	synthetic primer			
30	1227	SAUGUSCIC PLIMET			
	<400>	13			
	ggcccg	gg			8
_					
55	~07.0×	1.4			
	<210> <211>	1 4 8			
	<212>				
		synthetic primer			
60					
	<400>	14			
	ggcccc	cg			8

	<210>	15	
	<211>	19	
	<212>	DNA	
5	<213>	synthetic primer	
-			
	<400>	15	19
	ttttaa	tgcc ategeoteg	7.3
10			
10	<210>	16	
	<211>	20	
	<212>	DNA	
	<213>	synthetic primer	
15		•	
	<400>	16	
	cttcag	cagt ccaagccctg	20
20	<210>	17	
20	<211>	20	
		DNA	
		synthetic primer	
	72207	oynana aza przesa	
25	<400>	17	
	cctggc	gtta aactgctttc	20
	•		
	<210>	18	
20	<211>		
30	<212>		
	<213>	synthetic primer	
		-In-motor Francis	
	<400>	19	
35	ccatat	agtt tattaatata acac	24
	~21A>	10	
	<210>		
40	<211>		
40	<212> <213>	DNA synthetic primer	
	~~~~	straight brings	
	<400>	19	
	tatgtt	gcaa gtaggtgatc	20
45			
	<210>	20	
	<211>	32	
	<212>		
50	<213>		
50	<b>\Z13</b> /	synthetic primer	
	<400>	20	
	acgcgt	cgac gtgtgttaca amatggaccg am	32
~~			
55	Z21.05	25	
	<210>		
	<211>		
	<212>		
<b>.</b>	<b>\215&gt;</b>	synthetic primer	
60	<400>	21	
		teeg etggetggae aegeaagaag ca	32
	-2-23	bbb-rakar urdraakad re	34

	<210>	22	
	<211>	27	
	<212>	DNA	
5	<213>	synthetic primer	
,		•	
	<400>	22	
	cgaget	cocc ctccgctcca taccact	27
10			
	<210>	23	
	<211>		
	<212>		
	<213>	synthetic primer	
15	<.00×		
	<400>		31
	cgcgga	teeg etggetttaa agaaageagt t	31
20	<210>	24	
	<211>	20	
	<212>	DNA	
	<213>	synthetic primer	
25	<400>		20
	catgtg	acag atogaaggaa	20
	,		
	<210>	25	
30	<211>	21	
50	<212>	DNA	
	<213>	synthetic primer	
	<400>	0.5	
26	<400>	25 ttat totattoaga c	21
35	4.5.544	ctat tetatetaga t	
	<210>	26	
	<211>	6	
40	<212>		
	<213>	synthetic peptide	
	<220>		
		misc_feature	
45		() ()	
70	<223>		
		• •	
	4455		
~ ~	<400>	26	
50		u Xaa Gly His His	
	1	5	
	<210>	27	
55	<211>	5	
	<212>	PRT	
	<213>	synthetic peptide	
66	<400>	27	
60		g Arg His His	
	1	5	

```
- 18 -
    <210> 28
    <211> 5
    <212> PRT
    <213> synthetic peptide
    <400> 28
    His Val Ala His His
10
    <210> 29
    <211> 5
    <212> PRT
    <213> synthetic peptide
15
    <220>
    <221> misc_feature
    <222> ()..()
    <223> Xaa at position 2 is any amino acid; Xaa at position 3 is any amino acid
20
    <400> 29
    His Xaa Xaa His His
25
    <210> 30
    <211> 6
    <212> PRT
30
    <213> synthetic peptide
    <220>
    <221> misc_feature
    <222> ()..()
35
    <223> Kaa at position 2 is any amino acid; Kaa at position 3 is any amino acid;
    Xaa at posiation 4 is any amino acid
    <400> 30
40
    His Xaa Xaa Xaa His His
    ı
                   5
45
    <210> 31
    <211> 383
    <212> PRT
    <213> Glycine maxFAD2-2
50
    <400> 31
    Met Gly Ala Gly Gly Arg Thr Asp Val Pro Pro Ala Asn Arg Lys Sor
55
    Glu Val Asp Pro Leu Lys Arg Val Pro Phe Glu Lys Pro Gln Phe Ser
    Leu Ser Gln Ile Lys Lys Ala Ile Pro Pro His Cys Phe Gln Arg Ser
60
    Val Leu Arg Ser Phe Ser Tyr Val Val Tyr Asp Leu Thr Ile Ala Phe
                            55
```

<211> 383

Cys Leu Tyr Tyr Val Ala Thr His Tyr Phe His Leu Leu Pro Gly Pro Leu Ser Phe Arg Gly Met Ala Ile Tyr Trp Ala Val Gln Gly Cys Ile Leu Thr Gly Val Trp Val Ile Ala His Glu Cys Gly His His Ala Phe 10 Ser Asp Tyr Gln Leu Leu Asp Asp Ile Val Gly Leu Ile Leu His Ser Ala Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser His Arg Arg His 15 His Sor Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys Gln Lys Ser Cys Ile Lys Trp Tyr Ser Lys Tyr Leu Asn Asn Pro Pro 20 165 Gly Arg Val Leu Thr Leu Ala Val Thr Leu Thr Leu Gly Trp Pro Leu 25 Tyr Leu Ala Leu Asn Val Ser Gly Arg Pro Tyr Asp Arg Phe Ala Cys His Tyr Asp Pro Tyr Gly Pro Ile Tyr Ser Asp Arg Glu Arg Leu Gln 30 Ile Tyr Ile Ser Asp Ala Gly Val Leu Ala Val Val Tyr Gly Leu Phe Arg Leu Ala Met Ala Lys Gly Leu Ala Trp Val Val Cys Val Tyr Gly Val Pro Leu Leu Val Val Asn Gly Phe Leu Val Leu Ile Thr Phe Leu 265 40 Gln His Thr His Pro Ala Leu Pro His Tyr Thr Ser Ser Glu Trp Asp Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Asp Tyr Gly Ile Leu 45 295 Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Leu 310 315 Phe Ser Thr Met Pro His Tyr His Ala Met Glu Ala Thr Lys Ala Ile 50 Lys Fro Ile Leu Gly Glu Tyr Tyr Arg Phe Asp Glu Thr Pro Phe Val 55 Lys Ala Met Trp Arg Glu Ala Arg Glu Cys Ile Tyr Val Glu Pro Asp Gin Ser Thr Glu Ser Lys Gly Val Phe Trp Tyr Ash Ash Lys Leu 60 <210> 32

<212> PRT <213> Arabidopsis thaliana FAD2 <400> 32 5 Met Gly Ala Gly Gly Arg Met Pro Val Pro Thr Ser Ser Lys Lys Ser Glu Thr Asp Thr Thr Lys Arg Val Pro Cys Glu Lys Pro Pro Phe Ser 10 15 20 25

Val Gly Asp Leu Lys Lys Ala Ile Pro Pro His Cys Phe Lys Arg Ser Ile Pro Arg Ser Phe Ser Tyr Leu Ile Ser Asp Ile Ile Ile Ala Ser Cys Phe Tyr Tyr Val Ala Thr Asn Tyr Phe Ser Leu Leu Pro Gln Pro Leu Ser Tyr Leu Ala Trp Pro Leu Tyr Trp Ala Cys Gln Gly Cys Val Leu Thr Gly Ile Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Asp Tyr Gln Trp Leu Asp Asp Thr Val Gly Leu Ile Phe His Ser 120 Phe Leu Leu Val Pro Tyr Phe Ser Trp Lys Tyr Ser His Arg Arg His His Ser Asn Thr Gly Ser Leu Glu Arg Asp Glu Val Phe Val Pro Lys Cln Lys Ser Ala Ile Lys Trp Tyr Gly Lys Tyr Lou Asn Asn Pro Leu Gly Arg Ile Met Met Leu Thr Val Gln Phe Val Leu Gly Trp Pro Leu 40 Tyr Leu Ala Phe Asn Val Ser Gly Arg Pro Tyr Asp Gly Phe Ala Cys 200 His Phe Phe Pro Asn Ala Pro Ile Tyr Asn Asp Arg Clu Arg Leu Gln Ile Tyr Leu Ser Asp Ala Gly Ile Leu Ala Val Cys Phe Gly Leu Tyr 235 50 Arg Tyr Ala Ala Ala Gln Gly Met Ala Ser Met Ile Cys Leu Tyr Gly 55 280 Asn Lys Val Phe His Asn Ile Thr Asp Thr His Val Ala His His Lou

Val Pro Leu Leu Ile Val Asn Ala Phe Leu Val Leu Ile Thr Tyr Leu

Gln His Thr His Pro Ser Leu Pro His Tyr Asp Ser Ser Glu Trp Asp

Trp Leu Arg Gly Ala Leu Ala Thr Val Asp Arg Asp Tyr Gly Ile Leu

- 21 -

315 320 305 310 Phe Ser Thr Met Pro His Tyr Asn Ala Met Glu Ala Thr Lys Ala Ile Lys Pro Ile Leu Gly Asp Tyr Tyr Gln Phe Asp Gly Thr Pro Trp Tyr Val Ala Met Tyr Arg Glu Ala Lys Glu Cys Ile Tyr Val Glu Pro Asp 10 Arg Glu Gly Asp Lys Lys Gly Val Tyr Trp Tyr Asn Asn Lys Leu 375 15 <210> 33 <211> 387 <212> PRT <213> Glycine max FAD2-1 20 <400> 33 Met Gly Leu Ala Lys Glu Thr Thr Met Gly Gly Arg Gly Arg Val Ala Lys Val Glu Val Gln Gly Lys Lys Pro Leu Ser Arg Val Pro Asn Thr Lys Pro Pro Phe Thr Val Gly Gln Leu Lys Lys Ala Ile Pro Pro His 30 Cys Phe Gln Arg Ser Leu Leu Thr Ser Phe Ser Tyr Val Val Tyr Asp Leu Ser Phe Ala Phe Ile Phe Tyr Ile Ala Thr Thr Tyr Phe His Leu 35 Leu Pro Gln Pro Phe Ser Leu Ile Ala Trp Pro Ile Tyr Trp Val Leu 40 Gln Gly Cys Leu Leu Thr Gly Val Trp Val Ile Ala His Glu Cys Gly His His Ala Phe Ser Lys Tyr Gln Trp Val Asp Asp Val Val Gly Leu 45 Thr Leu His Ser Thr Leu Leu Val Pro Tyr Phe Ser Trp Lys Ile Ser 50 His Arg Arg His His Ser Asn Thr Gly Ser Leu Asp Arg Asp Glu Val Phe Val Pro Lys Pro Lys Ser Lys Val Ala Trp Phe Ser Lys Tyr Leu 55 Asn Asn Pro Leu Gly Arg Ala Val Ser Leu Leu Val Thr Leu Thr Ile Gly Trp Pro Met Tyr Leu Ala Phe Ash Val Ser Gly Arg Pro Tyr Asp 60 200 Ser Phe Ala Ser His Tyr His Pro Tyr Ala Pro Ile Tyr Ser Ash Arg

	Glu 225	Arg	Leu	Leu	Ile	Tyr 230	Val	Ser	yab	Val	Ala 235	Leu	Phe	Ser	Val	Thr 240
5	Tyr	Ser	Leu	Tyr	Arg 245	Val	Ala	Thr	Lou	1ys 250	Gly	Leu	Val	Trp	Leu 255	Leu
• •	СХг	Val	Tyr	Gly 260	Val	Pro	Leu	Leu	11e 265	Val	Asn	GLY	Phe	Leu 270	Val	Thr
10	Ile	Thr	<b>Tyr</b> 275	Leu	Gln	His	Thr	His 280	Phe	Ala	Leu	Pro	His 285	Tyr	yab	Ser
15	Ser	Glu 290	Trp	Asp	Trp	Leu	Lys 295	Gly	Ala	Lou	Ala	Thr 300	Met	Asp	Arg	Asp
	Tyr 305	Gly	Ile	Leu	Asn	Lys 310	Val	Phe	His	His	Ile 315	Thr	Asp	Thr	His	Val 320
20	Ala	His	His	Leu	Phe 325	Ser	Thr	Met	Pro	His 330	Tyr	His	Ala	Met	Glu 335	Ala
25	Thr	Asn	Ala	11e 340	ГĀв	Pro	Ile	Leu	Gly 345	Glu	Tyr	Tyr	Gln	Phe 350	Asp	Asp
ر ح	Thr	Pro	Phe 355	Tyr	Lys	Ala	Leu	Trp 360	Arg	Glu	Ala	Arg	<b>Glu</b> 365	Cys	Leu	Туг
30	Val	Glu 370	Pro	Азр	Glu	Gly	Thr 375	Ser	Glu	Lys	Gly	Val 380	Tyr	Trp	Tyr	Arg
	Asn 385	ГЛЗ	TYE													
35																
<b>4</b> 0	<210 <211 <212 <213	l> : 2> 1	34 377 PRT Brass	sica	napı	19 F/	<b>£</b> 04									
	<400	> :	34													
45	Met 1	Val	Val	Ala	Met 5	Asp	Gln	Arg	Ser	Asn 10	Ala	Asn	Gly	Авр	Glu 15	Arg
	Phe	Asp	Pro	Ser 20	Ala	Cln	Pro	Pro	Phe 25	L <b>y</b> s	Ile	Gly	Asp	I1⊕ 30	Arg	Ala
50	Ala	Ile	Pro 35	Lys	His	СХз	Trp	Val 40	Lys	Ser	Pro	Leu	Arg 45	Ser	Met	Ser
<b>&lt;</b> <	Tyr	Val 50	Ala	Arg	Asp	Ile	Phe 55	Ala	Val	Val	Ala	Leu 60	Ala	Val	Ala	Ala
55	Val 65	Tyr	Phe	Авр	Ser	Trp 70	Phe	Pho	ŢZP	Pro	Leu 75	Tyr	Trp	Ala	Ala	Gln BO
60	erĀ	Thr	Leu	Phe	Trp 85	Ala	Ile	Phe	Val	100 100	Gly	His	Asp	Суз	Gly 95	His
	Gly	Ser	Phe	Ser 100	Asp	Ile	Pro	Leu	Leu 105	naA	Thr	Ala	Val	Gly 110	His	Ile

			115					120			GIJ	~ <b>~</b> .	125		261	***
5	Arg	Thr 130	His	His	Gln	Asn	His 135	_	His	Val	Glu	Asn 140	Asp	G1u	Ser	Tr.
10	Val 145	Pro	Leu	Pro	Glu	Lys 150	Leu	Tyr	Lys	Asn	Leu 155	Ser	His	Ser	Thr	Ar 16
10	Met	Leu	Arg	Tyr	Thr 165	Val	Pro	Leu	Pro	Met 170	Leu	Ala	Tyr	Pro	Leu 175	Ty:
15	Leu	Trp	Tyr	Arg 180	Ser	Pro	ely	Lys	Glu 185	Gly	Ser	His	Tyr	Asn 190	Pro	Ty:
	Ser	Ser	Leu 195	Phe	Ala	Pro	Ser	<b>Gl</b> u 200	Arg	Lys	Leu	Ile	Ala 205	Thr	Ser	Thi
20	Thr	Cys 210	Trp	Ser	Ile	Met	Leu 215	Ala	Thr	Leu	Val	<b>Tyr</b> 220	Leu	Ser	Phe	Lei
25	Val 225	Gly	Pro	Val	Thr	Val 230	Leu	ŗ¥a	Val	Tyr	Gly 235	۷al	Pro	Tyr	Ile	11e 240
2.5	Phe	Val	Met	Trp	Leu 245	Asp	Ala	Val	Thr	Tyr 250	Leu	His	His	His	Gly 255	
30	Asp	Asp	Lys	Leu 260	Pro	Trp	TYE	Arg	G1 <b>y</b> 265	Lys	Glu	Trp	Ser	<b>Tyr</b> 270	Leu	Arg
	61¥	еĵÃ	Leu 275	Thr	Thr	Ile	Авр	Arg 280	Asp	Tyr	Gly	Ile	Phe 285	Asn	neA	Ile
35	His	His 290	Asp	Ile	C1Y	Thr	His 295	Val	Ile	His	His	<b>Le</b> u 300	Phe	Pro	Gln	Ile
40	Pro 305	His	Tyr	His	Leu	Val 310	Yab	Ala	Thr	Lys	Ser 315	Ala	Lys	Ris	Val	Lev 320
	Gly	Arg	Tyr	Tyr	Arg 325	G1u	Pro	ГÃЗ	Thr	Ser 330	ĠŢĀ	Ala	Ile	Pro	Ile 335	His
45	Leu	Val	Glu	Ser 340	Leu	Val	Ala	Ser	11e 345	Lys	Lys	Asp	His	<b>Tyr</b> 350	Val	Sez
			Gly 355					360		Thr	Asp	Pro	Аяр 365	Leu	Tyr	Va)
50	Tyr	Ala 370	Ser	Asp	ГÀв	Ser	Lys 375	Ile	ABN							
55	<210 <211 <212 <213	.> 4 !> E	35 124 PRT Slyci	.nema	LX F7	MD6										
60	<400		35			-										
00	Met 1	Ala	Cys	Thr	Leu 5	Ala	Asp	Ser	Leu	Leu 10	Leu	Phe	Lys	Gly	Ser 15	Τχι

Gln Lys Pro Val Leu Arg Arg Asp Ile Ala Ala Arg Tyr Ser Pro Gly 20 25 Ile Phe Ser Leu Asn Ser Asn Gly Leu Ile Gln Lys Arg Phe Arg Arg 5 Gln Arg Asn Phe Val Thr Arg Asn Lys Val Thr Val Ile His Ala Val Ala Ile Pro Val Gln Pro Ala Pro Val Glu Ser Ala Glu Tyr Arg Lys 10 Cln Leu Ala Glu Asp Tyr Gly Phe Arg Gln Val Gly Glu Pro Leu Ser 15 Asp Asp Val Thr Leu Lys Asp Val Ile Asm Pro Leu Pro Lys Glu Val 105 Phe Glu Ile Asp Asp Val Lys Ala Trp Lys Ser Val Leu Ile Ser Val 20 Thr Ser Tyr Ala Leu Gly Leu Phe Met Ile Ser Lys Ala Pro Trp Tyr Leu Lou Pro Leu Ala Trp Val Trp Thr Gly Thr Ala Ile Thr Gly Phe 25 Phe Val Ile Gly His Asp Cys Ala His Arg Ser Phe Ser Ser Asn Lys 30 Leu Val Glu Asp Ile Val Gly Thr Leu Ala Phe Met Pro Leu Ile Tyr Pro Tyr Glu Pro Trp Arg Phe Lys His Asp Arg His His Ala Lys Thr 35 200 Asn Met Leu Arg Glu Asp Thr Ala Trp His Pro Val Trp Lys Asp Glu 40 Phe Glu Ser Thr Pro Leu Leu Arg Lys Ala Ile Ile Tyr Gly Tyr Gly Pro Phe Arg Cys Trp Met Ser Ile Ala His Trp Leu Met Trp His Phe 245 45 Asp Leu Lys Lys Phe Arg Pro Ser Glu Val Pro Arg Val Lys Ile Ser Leu Ala Cys Val Phe Ala Phe Ile Ala Ile Gly Trp Pro Leu Ile Ile 50 280 Tyr Lys Thr Gly Ile Met Gly Trp Ile Lys Phe Trp Leu Met Pro Trp Lou Cly Tyr His Pho Trp Met Ser Thr Phe Thr Met Val His His Thr 55 310 Ala Pro Tyr Ile Pro Phe Lys Tyr Ser Glu Glu Trp Asn Arg Ala Gln 60 Ala Gln Leu Asn Gly Thr Val His Cys Asp Tyr Pro Lys Trp Ile Glu

Ile	Leu	Cy≄ 355	His	Asp	Ile	Asn	Val 360	His	Ile	Pro	His	His 365	Ile	Ser	Pr
Arg	Ile 370	Pro	Ser	Tyr	Asn	<b>Leu</b> 375	Arg	Ala	Ala	His	<b>Lys</b> 380	Ser	Leu	Gln	Gli
Asn 385	Trp	Gly	Gln	Tyr	Leu 390	neA	Glu	Ala	Ser	Ттр 395	Asn	Trp	Arg	Leu	Me:
Lys	Thr	Ile	Met	Thr 405	Val	Сув	Gln	Val	Tyr 410	Asp	Lys	Glu	Lys	Ser 415	Lei
Суз	Сув	Leu	Arg	Arg	Thr	Cvs	Pro								

Cys Cys Leu Arg Arg Thr Cys Pr 420

15

10